



SEQUENCE LISTING

KADOWAKI, Takashi  
YAMAUCHI, Toshimasa  
NAGAI, Ryoza  
KAMON, Jyunji

<120> ADIPONECTIN RECEPTOR AND GENE ENCODING  
THE SAME

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<140> 10/799943

<141> 2004-03-11

<150> PCT/JP03/07515

<151> 2003-06-12

<150> JP 2002-383738

<151> 2002-12-29

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Ala Ser Asn Arg Glu Ala Asp Thr Val Glu Leu Ala Glu Leu Gly Pro  
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ctg cta gaa gag aag ggc aaa cgg gta atc gcc aac cca ccc aaa gct 144  
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35 40 45

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Glu Glu Glu Gln Thr Cys Pro Val Pro Gln Glu Glu Glu Glu Val  
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cgg gta ctg aca ctt ccc ctg caa gcc cac cac gcc atg gag aag atg 240  
Arg Val Leu Thr Leu Pro Leu Gln Ala His His Ala Met Glu Lys Met  
65 70 75 80

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Tyr Gly Val Ser Asn Leu Gln Glu Phe Arg Tyr Gly Leu Glu Gly Gly				
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Gly His Arg Pro Pro Met Pro Ser Phe Arg Ala Cys Phe Lys Ser Ile																
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 100 105 110  
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Trp	Phe	His	Ser	His	Gln	Leu	Phe	His	Ile	Phe	Val	Val	Ala	Gly	Ala	
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Tyr	Val	Ile	Ser	Glu	Gly	Phe	Leu	Lys	Ala	Ala	Thr	Ile	Gly	Gln	Ile		
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Tyr	Ala	Ala	Arg	Ile	Pro	Glu	Arg	Phe	Phe	Pro	Gly	Lys	Cys	Asp	Ile		
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Ser	Gly	Asn	Arg	Glu	Ala	Asp	Thr	Val	Glu	Leu	Ala	Glu	Leu	Gly	Pro	
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Tyr	Tyr	Ser	Phe	Tyr	Cys	Ser	Pro	Gln	Pro	Arg	Leu	Ile	Tyr	Leu	Ser		
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His	Gln	Ile	Phe	His	Val	Leu	Val	Val	Ala	Ala	Ala	Phe	Val	His	Phe			
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tat	ggg	gtg	tcc	aac	ctt	cag	gaa	ttc	cgt	tat	ggc	cta	gaa	ggg	ggc	1104		
Tyr	Gly	Val	Ser	Asn	Leu	Gln	Glu	Phe	Arg	Tyr	Gly	Leu	Glu	Gly	Gly			
355				360				365										
tgt	acc	gac	gac	tcc	ctt	ctc	tga								1128			
Cys	Thr	Asp	Asp	Ser	Leu	Leu												
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 35 40 45  
 Glu Glu Asp Gln Ala Cys Pro Val Pro Gln Glu Glu Glu Glu Val  
 50 55 60  
 Arg Val Leu Thr Leu Pro Leu Gln Ala His His Ala Met Glu Lys Met  
 65 70 75 80  
 Glu Glu Phe Val Tyr Lys Val Trp Glu Gly Arg Trp Arg Val Ile Pro  
 85 90 95  
 Tyr Asp Val Leu Pro Asp Trp Leu Lys Asp Asn Asp Tyr Leu Leu His  
 100 105 110  
 Gly His Arg Pro Pro Met Pro Ser Phe Arg Ala Cys Phe Lys Ser Ile  
 115 120 125  
 Phe Arg Ile His Thr Glu Thr Gly Asn Ile Trp Thr His Leu Leu Gly  
 130 135 140  
 Phe Val Leu Phe Leu Phe Leu Gly Ile Leu Thr Met Leu Arg Pro Asn



145		150		155		160
Met Tyr Phe Met	Ala Pro Leu Gln Glu Lys Val Val Phe Gly Met Phe					
	165	170	175			
Phe Leu Gly Ala Val Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr						
	180	185	190			
Val Tyr Cys His Ser Glu Lys Val Ser Arg Thr Phe Ser Lys Leu Asp						
	195	200	205			
Tyr Ser Gly Ile Ala Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu						
	210	215	220			
Tyr Tyr Ser Phe Tyr Cys Ser Pro Gln Pro Arg Leu Ile Tyr Leu Ser						
	225	230	235			
Ile Val Cys Val Leu Gly Ile Ser Ala Ile Ile Val Ala Gln Trp Asp						
	245	250	255			
Arg Phe Ala Thr Pro Lys His Arg Gln Thr Arg Ala Gly Val Phe Leu						
	260	265	270			
Gly Leu Gly Leu Ser Gly Val Val Pro Thr Met His Phe Thr Ile Ala						
	275	280	285			
Glu Gly Phe Val Lys Ala Thr Thr Val Gly Gln Met Gly Trp Phe Phe						
	290	295	300			
Leu Met Ala Val Met Tyr Ile Thr Gly Ala Gly Leu Tyr Ala Ala Arg						
	305	310	315			
Ile Pro Glu Arg Phe Phe Pro Gly Lys Phe Asp Ile Trp Phe Gln Ser						
	325	330	335			
His Gln Ile Phe His Val Leu Val Val Ala Ala Ala Phe Val His Phe						
	340	345	350			
Tyr Gly Val Ser Asn Leu Gln Glu Phe Arg Tyr Gly Leu Glu Gly Gly						
	355	360	365			
Cys Thr Asp Asp Ser Leu Leu						
	370	375				

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Glu Glu Phe Val Cys Lys Val Trp Glu Gly Arg Trp Arg Val Ile Pro	
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cac gat gtg cta ccg gat tgg ctt aag gat aat gac ttc ctt ctc cat	144
His Asp Val Leu Pro Asp Trp Leu Lys Asp Asn Asp Phe Leu Leu His	
35 40 45	
gga cac cgg cct cct atg cct tcc ttt cgg gcc tgt ttt aag agc att	192
Gly His Arg Pro Pro Met Pro Ser Phe Arg Ala Cys Phe Lys Ser Ile	
50 55 60	

ttt aga ata cac aca gag acg ggc aac att tgg aca cat ctc cta ggt	240
Phe Arg Ile His Thr Glu Thr Gly Asn Ile Trp Thr His Leu Leu Gly	
65 70 75 80	
tgt gta ttc ttc ctg tgc ctg ggg atc ttt tat atg ttt cgc cca aat	288
Cys Val Phe Phe Leu Cys Leu Gly Ile Phe Tyr Met Phe Arg Pro Asn	
85 90 95	
ata tct ttt gtg gcc cct ctg caa gag aaa gtg gtc ttt ggc ttg ttc	336
Ile Ser Phe Val Ala Pro Leu Gln Glu Lys Val Val Phe Gly Leu Phe	
100 105 110	
ttc ttg gga gcc att ctc tgc ctt tcc ttt tca tgg ctc ttc cac acg	384
Phe Leu Gly Ala Ile Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr	
115 120 125	
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Val Tyr Cys His Ser Glu Gly Val Ser Arg Leu Phe Ser Lys Leu Asp	
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Tyr Ser Gly Ile Ala Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu	
145 150 155 160	
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Tyr Tyr Ser Phe Tyr Cys Asn Pro Gln Pro Cys Phe Ile Tyr Leu Ile	
165 170 175	
gtc atc tgt gtg ctg ggc att gca gcc att atc gtc tct cag tgg gac	576
Val Ile Cys Val Leu Gly Ile Ala Ala Ile Ile Val Ser Gln Trp Asp	
180 185 190	
atg ttt gcc acc cct cag tat cgg ggg gtc aga gca gga gtg ttc gtg	624
Met Phe Ala Thr Pro Gln Tyr Arg Gly Val Arg Ala Gly Val Phe Val	
195 200 205	
ggc tta ggc ctg agt gga atc atc cct acc ttg cat tat gtc atc tca	672
Gly Leu Gly Leu Ser Gly Ile Ile Pro Thr Leu His Tyr Val Ile Ser	
210 215 220	
gaa ggg ttc ctg aag gct gcc acc ata ggg cag ata ggc tgg cta atg	720
Glu Gly Phe Leu Lys Ala Ala Thr Ile Gly Gln Ile Gly Trp Leu Met	
225 230 235 240	
ctt atg gct agc ctc tat atc acc gga gct gcc ctc tat gcg gcc cgt	768
Leu Met Ala Ser Leu Tyr Ile Thr Gly Ala Ala Leu Tyr Ala Ala Arg	
245 250 255	
atc cct gag cgc ttc ttt cct ggc aaa tgt gac atc tgg ttt cac tct	816
Ile Pro Glu Arg Phe Phe Pro Gly Lys Cys Asp Ile Trp Phe His Ser	
260 265 270	
cat cag ctc ttc cac atc ttt gtg gtt gct ggt gcc ttt gtt cac ttc	864
His Gln Leu Phe His Ile Phe Val Val Ala Gly Ala Phe Val His Phe	
275 280 285	
cac gga gtc tca aac ctg cag gaa ttt cgt ttc atg att ggc ggg ggc	912

His Gly Val Ser Asn Leu Gln Glu Phe Arg Phe Met Ile Gly Gly Gly  
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936

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 35 40 45  
 Gly His Arg Pro Pro Met Pro Ser Phe Arg Ala Cys Phe Lys Ser Ile  
 50 55 60  
 Phe Arg Ile His Thr Glu Thr Gly Asn Ile Trp Thr His Leu Leu Gly  
 65 70 75 80  
 Cys Val Phe Phe Leu Cys Leu Gly Ile Phe Tyr Met Phe Arg Pro Asn  
 85 90 95  
 Ile Ser Phe Val Ala Pro Leu Gln Glu Lys Val Val Phe Gly Leu Phe  
 100 105 110  
 Phe Leu Gly Ala Ile Leu Cys Leu Ser Phe Ser Trp Leu Phe His Thr  
 115 120 125  
 Val Tyr Cys His Ser Glu Gly Val Ser Arg Leu Phe Ser Lys Leu Asp  
 130 135 140  
 Tyr Ser Gly Ile Ala Leu Leu Ile Met Gly Ser Phe Val Pro Trp Leu  
 145 150 155 160  
 Tyr Tyr Ser Phe Tyr Cys Asn Pro Gln Pro Cys Phe Ile Tyr Leu Ile  
 165 170 175  
 Val Ile Cys Val Leu Gly Ile Ala Ala Ile Ile Val Ser Gln Trp Asp  
 180 185 190  
 Met Phe Ala Thr Pro Gln Tyr Arg Gly Val Arg Ala Gly Val Phe Val  
 195 200 205  
 Gly Leu Gly Leu Ser Gly Ile Ile Pro Thr Leu His Tyr Val Ile Ser  
 210 215 220  
 Glu Gly Phe Leu Lys Ala Ala Thr Ile Gly Gln Ile Gly Trp Leu Met  
 225 230 235 240  
 Leu Met Ala Ser Leu Tyr Ile Thr Gly Ala Ala Leu Tyr Ala Ala Arg  
 245 250 255  
 Ile Pro Glu Arg Phe Phe Pro Gly Lys Cys Asp Ile Trp Phe His Ser  
 260 265 270  
 His Gln Leu Phe His Ile Phe Val Val Ala Gly Ala Phe Val His Phe  
 275 280 285  
 His Gly Val Ser Asn Leu Gln Glu Phe Arg Phe Met Ile Gly Gly Gly  
 290 295 300  
 Cys Thr Glu Glu Asp Ala Leu  
 305 310